

WO 99/51771

09/647596

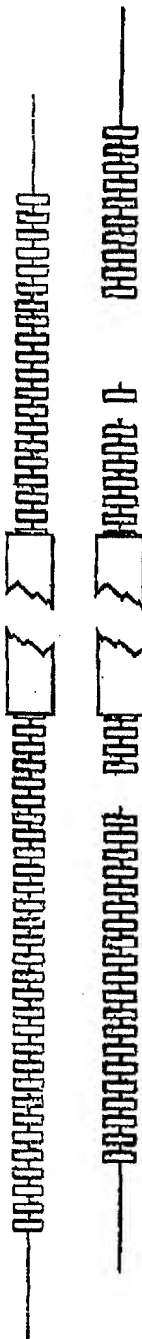
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Fig 1

M.bovis BCG

IS6110



M.tuberculosis H37Rv

Consensus DR sequence: GTCGTCAGACCCAAACCCCGAGAGGGACGGAAC



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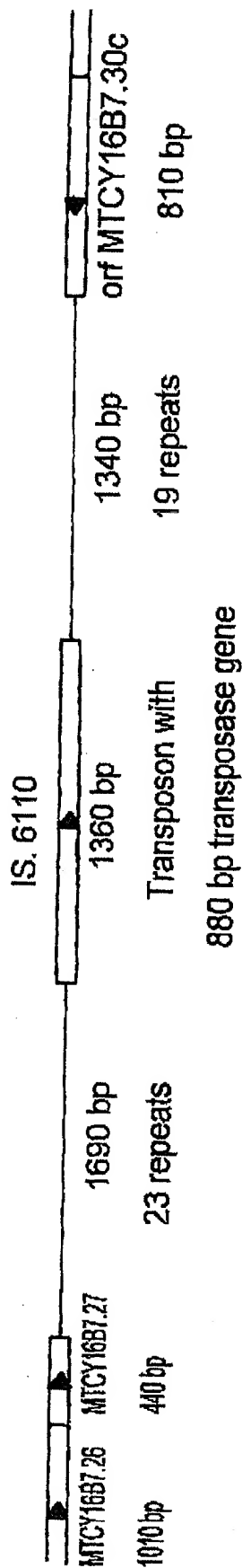
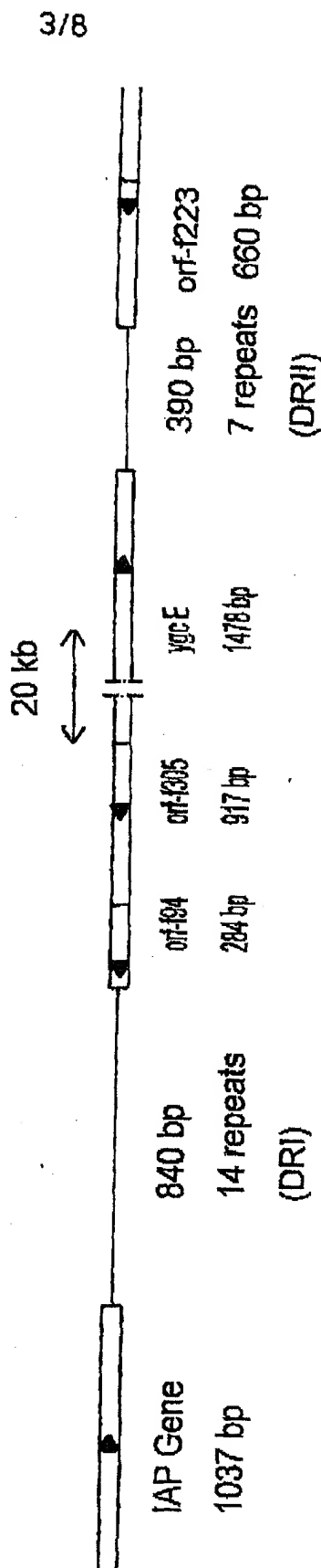
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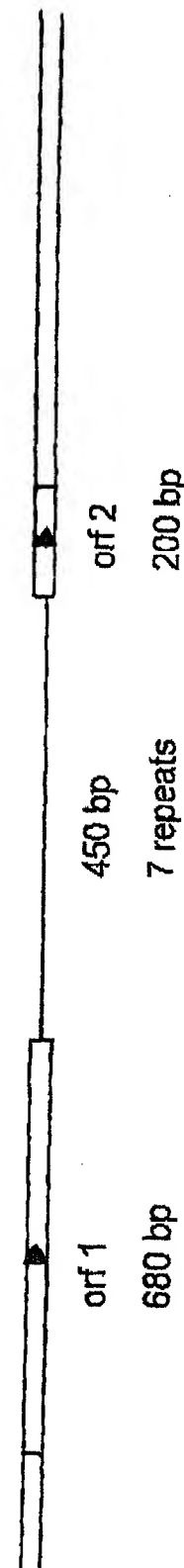
Fig 3.1

## M. TUBERCULOSIS

H37Rv

ESCHERICHIA COLI  
K12

## STREPTOCOCCUS PYOGENES



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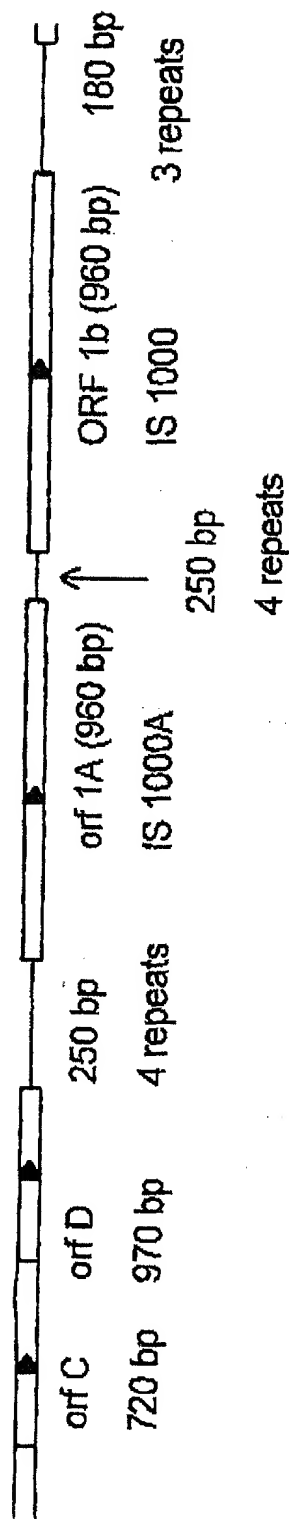
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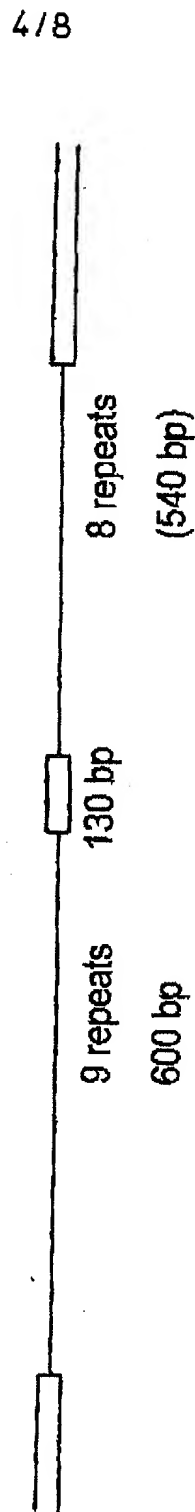
## THERMUS AQUATICUS THERMOPHILUS

HB8 ATCC 27634

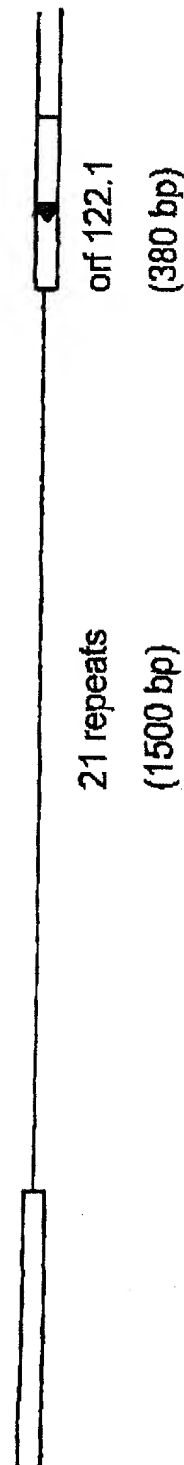
Fig 3.2



## ANABAENA sp. PCC.7120



## HALOFERAX MEDITERRANEI



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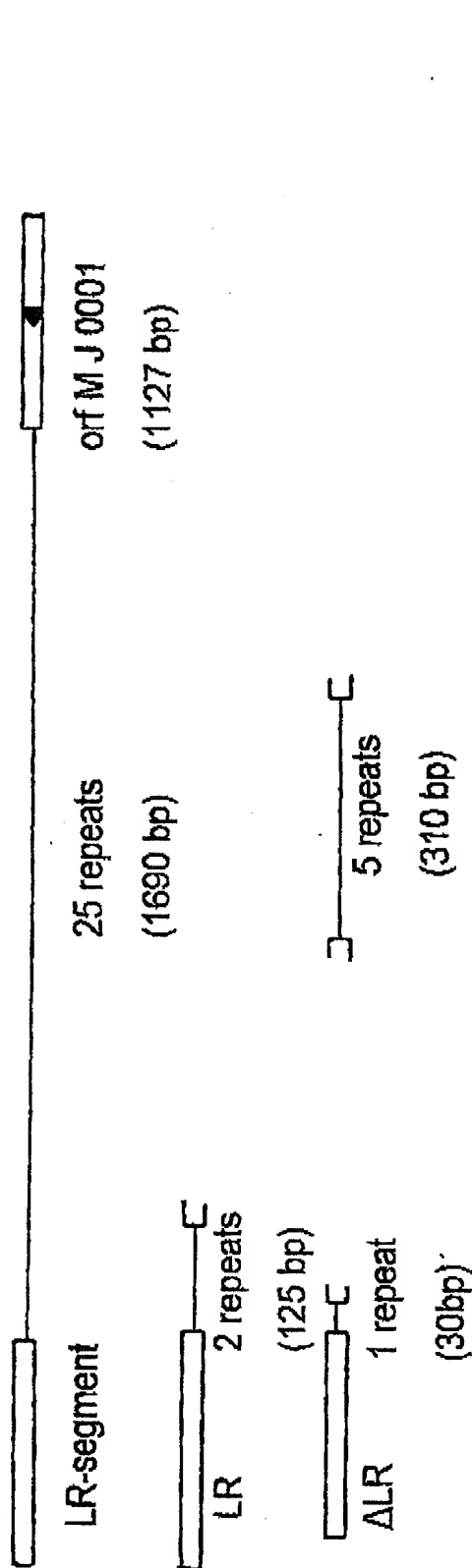
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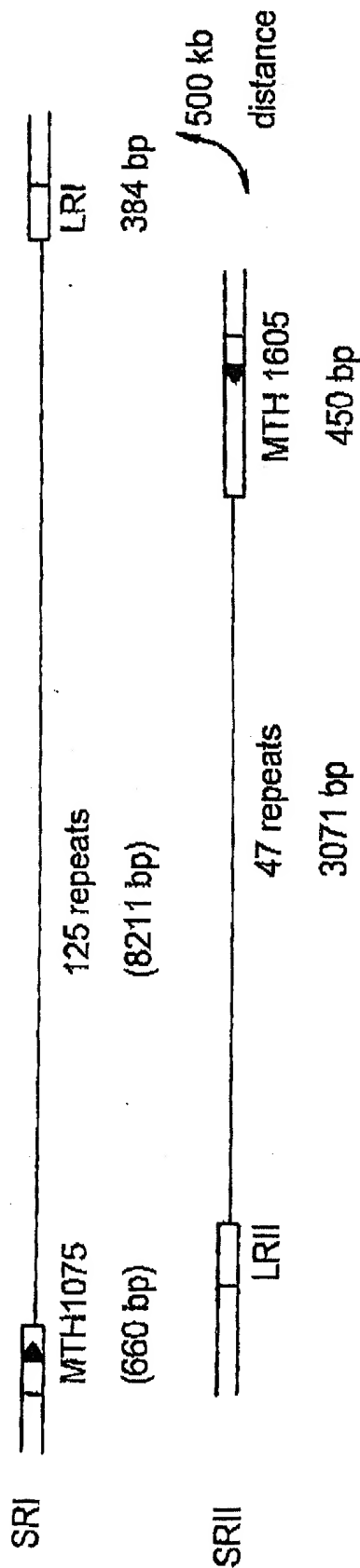
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# METHANOCOCCUS JANNASCHII

Fig 3.3



# METHANOBACTERIUM THERMOAUTOTROPHICUM



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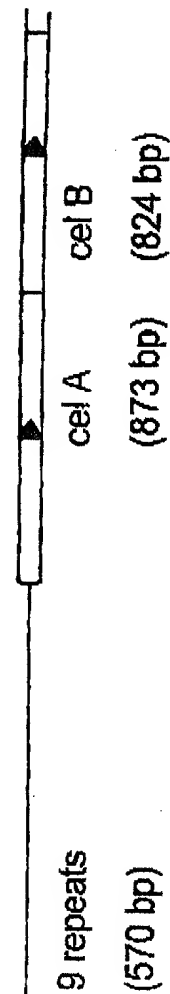
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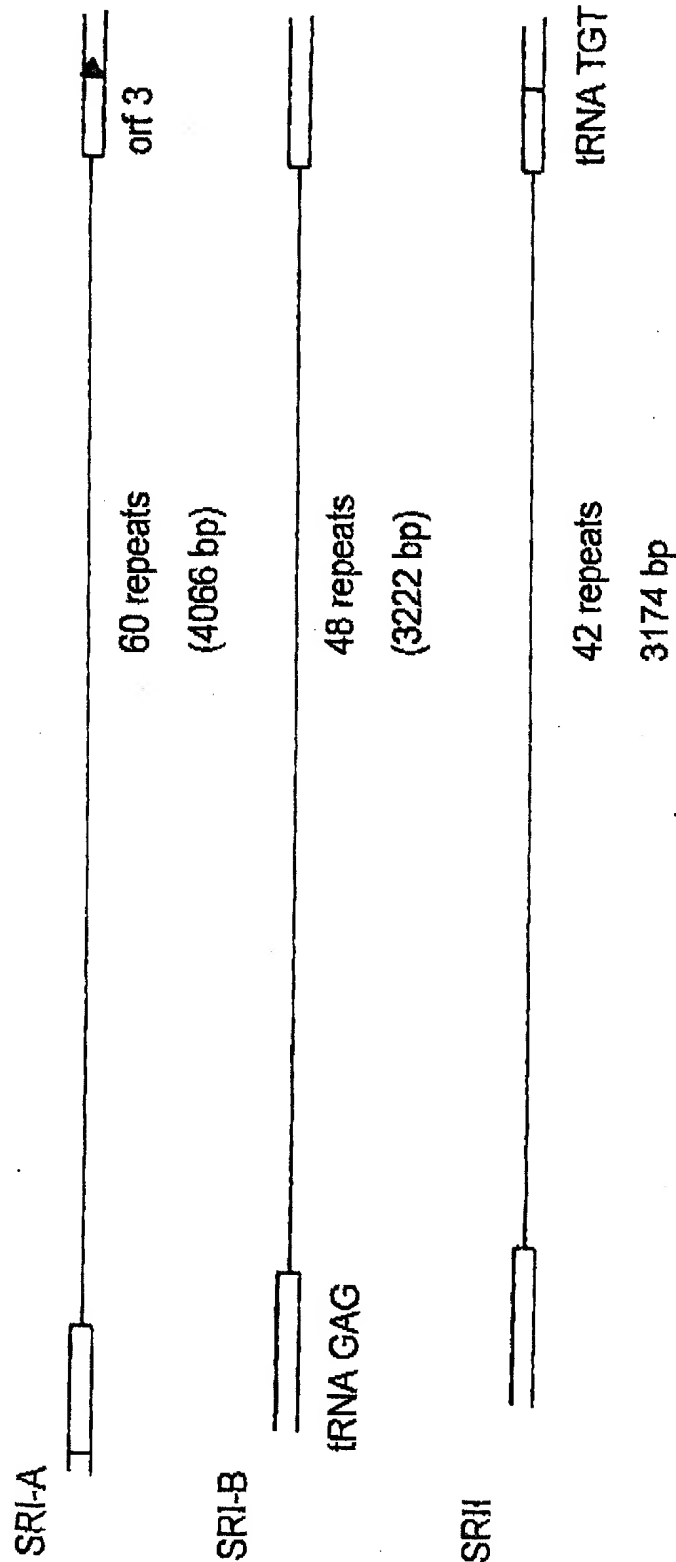
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THERMOTOGA MARITIMA

Fig 3<sup>4</sup>



ARCHAEOGLOBUS FULGIDUS



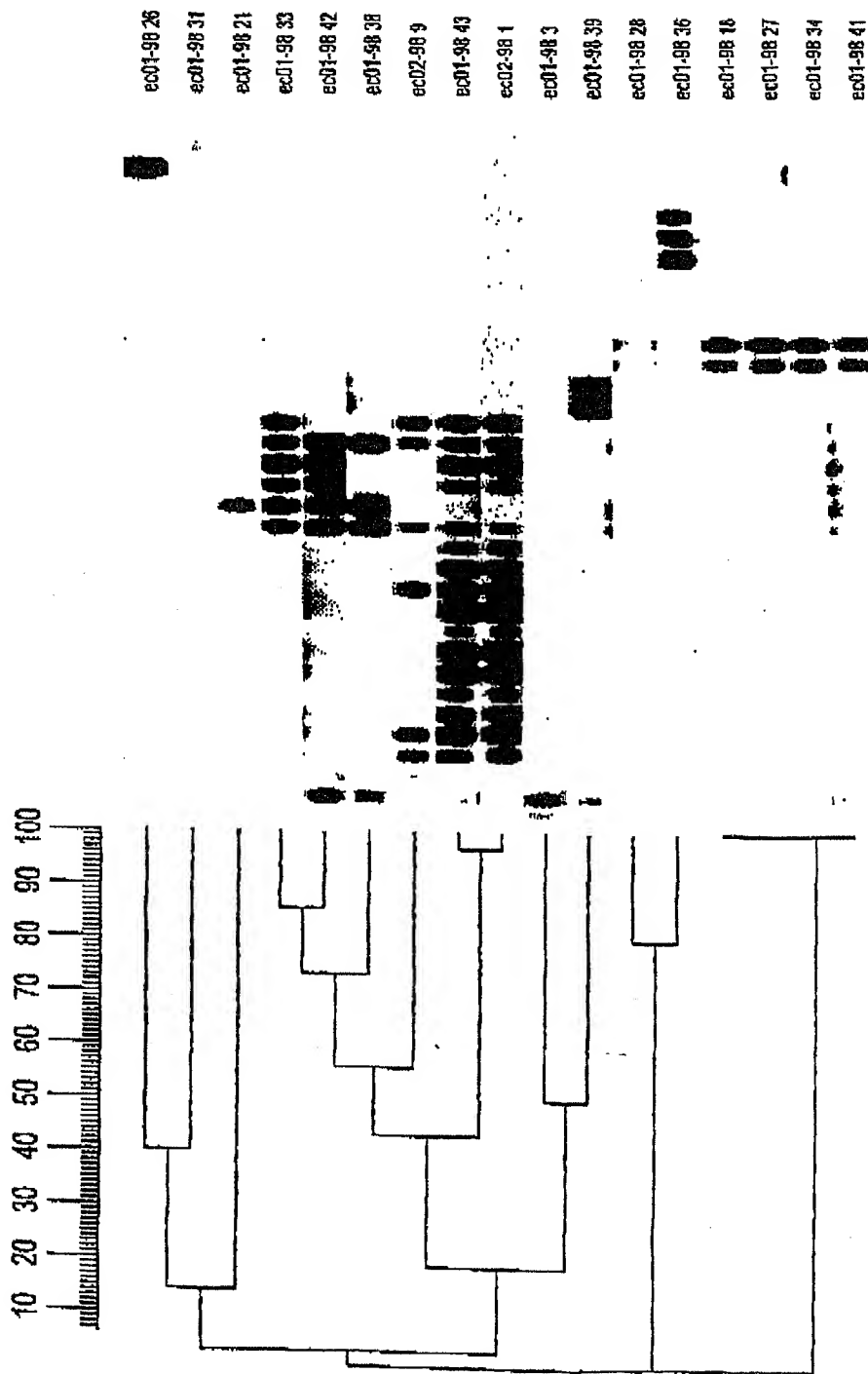
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Fig 4

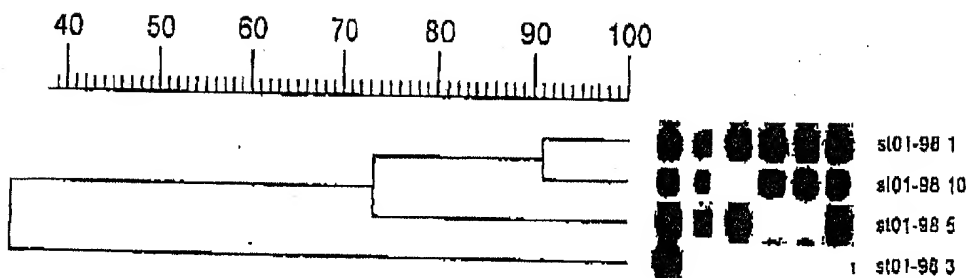


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*Fig 5*





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**Fig. 6**

SEQ ID No. 1: GTCGTCAGACCCAAAACCCCGAGAGGGGACGGAAAC

SEQ ID No. 2: CGGTTTATCCCCGCTGGCGCGGG<sup>G</sup>AACTC

SEQ ID No. 3: CGGTTTATCCCCGCTGGCGCGGGGAACTC

SEQ ID No. 4: CGGTTTATCCCCGCTGGCGCGGGGAACTC

SEQ ID No. 5: CGGTTTATCCCCGCTGGCGCGGGGAACTC

SEQ ID No. 6: CGGTTTATCCCCGCTGGCGCGGGGAACTC

SEQ ID No. 7: CGGTTTATCCCCGCTGGCGCGGGGAACTC

SEQ ID No. 8: CGGTTTATCCCCGCTGGCGCGGATACAC

SEQ ID No. 9: GTTTTAGAGCTATGCTGTTTTGAATGGTCCCAAAC

SEQ ID No. 10: AATCCCCTTACGGGGCTCAATCCCTTGCAA

SEQ ID No. 11: GTTTCATACTTCCTTAGAGGTATGGAAAC

SEQ ID No. 12: GTTTTAACTAACA AAAATCCCTATCAGGGATTGAAAC

SEQ ID No. 13: GTTTAAACTTTATAAAAATCCCTTTTAGGGATTGAAAC

SEQ ID No. 14: GTTACAGACGAACCCTAGTTGGGTTGAAGC

SEQ ID No. 15: AATTAAAATCAGACCGTTTCGGAATGGAAA

SEQ ID No. 16: ATTTCAATCCCATTGTTGGTCTGATTTTAAC

SEQ ID No. 17: GTTAAAATCAGACCAAATGGGATTGAAAT

SEQ ID No. 18: CTTTCAATCCCATTGTTGGTCTGATTTCAAC

AMENDED SHEET

**Fig. 6**

SEQ ID No. 1: GTCGTCAGACCCAAAACCCCGAGAGGGGACGGAAAC  
SEQ ID No. 2: CGGTTTATCCCCGCTGGCGCGGGAACTC  
SEQ ID No. 3: CGGTTTATCCCCGCTGGCGCGGGGAACTC  
SEQ ID No. 4: CGGTTTATCCCCGCTGGCGCGGGGAACTC  
SEQ ID No. 5: CGGTTTATCCCCGCTGGCGCGGGGAACTC  
SEQ ID No. 6: CGGTTTATCCCCGCTGGCGCGGGGAACTC  
SEQ ID No. 7: CGGTTTATCCCCGCTGGCGCGGGGAACTC  
SEQ ID No. 8: CGGTTTATCCCCGCTGGCGCGGATACAC  
SEQ ID No. 9: GTTTTAGAGCTATGCTGTTTTGAATGGTCCCAAAC  
SEQ ID No. 10: AATCCCCTTACGGGGCTCAATCCCTTGCAA  
SEQ ID No. 11: GTTTCATACTTCCTTAGAGGTATGGAAAC  
SEQ ID No. 12: GTTTTAACTAACAAAAATCCCTATCAGGGATTGAAAC  
SEQ ID No. 13: GTTTAACTTTATAAAATCCCTTTTAGGGATTGAAAC  
SEQ ID No. 14: GTTACAGACGAACCCTAGTTGGGTTGAAGC  
SEQ ID No. 15: AATTAAAATCAGACCGTTTCGGAATGGAAA  
SEQ ID No. 16: ATTTCAATCCCATTTTGGTCTGATTTTAAC  
SEQ ID No. 17: GTTAAAATCAGACCAAATGGGATTGAAAT  
SEQ ID No. 18: CTTTCAATCCCATTTTGGTCTGATTTCAAC